Claim 43, line 3, after "pCR1" change the comma "," to --and--, and delete "and pSC1".

Claim 44, line 4, delete "at least".

Claim 45, line 4, delete "at least";

line 9 of the sequence, change "Ais" to --

## Please add the following new Claims 49 to 59.

- --49. A recombinant expression plasmid for producing human fibroblast  $\beta_1$  interferon, comprising a DNA sequence coding for human fibroblast  $\beta_1$  interferon, whereby expression of human fibroblast  $\beta_1$  interferon by a host is enabled.
- 50. A recombinant expression plasmid for producing human fibroblast  $\beta_1$  interferon polypeptide in a host, comprising a DNA sequence coding for human fibroblast  $\beta_1$  interferon operably linked therein for expression.

fibroblast  $\beta_1$  interferon polypeptide, transformed with an expression plasmid of claim  $\frac{1}{39}$ .

interferon polypeptide, comprising expressing the expression plasmid of claim 49 in a host cell.

53. A human fibroblast  $\beta_1$  interferon polypeptide produced by the process of claim 52.

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- 54. Recombinant human Fibroblast  $\beta_1$  interferon.
- 31 55. A plasmid of claim 49, wherein said DNA has the

## sequence:

(3/0X

ATG ACC AAC AAG TGT CTC CTC CAA ATT GCT CTC CTG TTG TGC TTC TCC TAC TGG TTG TTC ACA GAG GAG GTT TAA CGA GAG GAC AAC ACG AAG AGG ACT ACA GCT CTT TCC ATG AGC TAC AAC TTG CTT GGA TTC CTA CAA AGA TGA TGT CGA GAA AGG TAC TCG ATG TTG AAC GAA CCT AAG CAT GTT TCT AGC AGC AAT TTT CAG TGT CAG AAG CTC CTG TGG CAA TTG AAT GGG AGG TCG TCG TTA AAA GTC ACA GTC TTC GAG GAC ACC GTT AAC TTA CCC TCC CTT GAA TAT TGC CTC AAG GAC AGG ATG AAC TTT GAC ATC CCT GAG GAG GAA CTT ATA ACG GAG TTC CTG TCC TAC TTG AAA CTG TAG GGA CTC CTC ATT AAG CAG CTG CAG CAG TTC CAG AAG GAC GCC GCA TTG ACC ATC TAA TTC GTC GAC GTC GTC AAG GTC TTC CTC CTG CGG CGT AAC TGG TAG TAT GAG ATG CTC CAG AAC ATC TTT GCT ATT TTC AGA CAA GAT TCA TCT ATA CTC TAC GAG GTC TTG TAG AAA CGA TAA AAG TCT GTT CTA AGT AGA AGC ACT GGC TGG AAT GAG ACT ATT GTT GAG AAC CTC CTG GCT AAT GTC TCG TGA CCG ACC TTA CTC TGA TAA CAA CTC TTG GAG GAC CGA TTA CAG TAT CAT CAG ATA AAC CAT CTG AAG ACA GTC CTG GAA GAA AAA CTG GAG ATA GTA GTC TAT TTG GTA GAC TTC TGT CAG GAC CTT CTT TTT GAC CTC AAA GAA GAT TTC ACC AGG GGA AAA CTC ATG AGC AGT CTG CAC CTG AAA TTT CTT CTA AAG TGG TCC CCT TTT GAG TAC TCG TCA GAC GTG GAC TTT AGA TAT TAT GGG AGG ATT CTG CAT TAC CTG AAG GCC AAG GAG TAC AGT TCT ATA ATA CCC TCC TAA GAC GTA ATG GAC TTC CGG TTC CTC ATG TCA CAC TGT GCC TGG ACC ATA GTC AGA GTG GAA ATC CTA AGG AAC TTT TAC GTG ACA CGG ACC TGG TAT CAG TCT CAC CTT TAG GAT TCC TTG AAA ATG TTC ATT AAC AGA CTT ACA GGT TAC CTC CGA AAC AAG TAA TTG TCT GAA TGT CCA ATG GAG GCT TTG.

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56.78 An Extensión cell hart 127, wherein said DNA has the

## sequence:

T311X

ATG ACC AAC AAG TGT CTC CTC CAA ATT GCT CTC CTG TTG TGC TTC TCC TAC TGG TTG TTC ACA GAG GAG GTT TAA CGA GAG GAC AAC ACG AAG AGG

ACT ACA GCT CTT TCC ATG AGC TAC AAC TTG CTT GGA TTC CTA CAA AGA TGA TGT CGA GAA AGG TAC TCG ATG TTG AAC GAA CCT AAG ØAT GTT TCT AGC AGC AAT TTT CAG TGT CAG AAG CTC CTG TGG CAA TTG AAT GGG AGG TCG TCG TTA AAA GTC ACA GTC TTC GAG GAC ACC GTT AAC TTA CCC TCC CTT GAA TAT TGC CTC AAG GAC AGG ATG AAC TTT GAC ATC CCT GAG GAG GAA CTT ATA ACG GAG TTC CTG TCC TAC TTG AAA CTG TAG GGA CTC CTC ATT AAG CAG CTG CAG CAG TTC CAG AAG GAG GAC GCC GCA TTG ACC ATC TAA TTC GTC GAC GTC AAG GTC TTC CTC CTG CGG CGT AAC TGG TAG TAT GAG ATG CTC CAG AAC ATC TTT GCT ATT TTC AGA CAA GAT TCA TCT ATA CTC TAC GAG GTC TTG TAG AAA CGA TAA AAG TCT GTT CTA AGT AGA AGC ACT GGC TGG AAT GAG ACT ATT GTT GAG AAC CTC CTG GCT AAT GTC TCG TGA CCG ACC TTA CTC TGA TAA CAA CTC TTG GAG GAC CGA TTA CAG TAT CAT CAG ATA AAC CAT CTG AAG ACA GTC CTG GAA GAA AAA CTG GAG ATA GTA GTC TAT TTG GTA GAC TTC TGT CAG GAC CTT CTT TTT GAC CTC AAA GAA GAT TTC ACC AGG GGA AAA CTC ATG AGC AGT CTG CAC CTG AAA TTT CTT CTA AAG TGG TCC CCT TTT GAG TAC TCG TCA GAC GTG GAC TTT AGA TAT TAT GGG AGG ATT CTG CAT TAC CTG AAG GCC AAG GAG TAC AGT TCT ATA ATA CCC TCC TAA GAC GTA ATG GAC TTC CGG TTC CTC ATG TCA CAC TGT GCC TGG ACC ATA GTC AGA GTG GAA ATC CTA AGG AAC TTT TAC GTG ACA CGG ACC TGG TAT CAG TCT CAC CTT TAG GAT TCC TTG AAA ATG TTC ATT AAC AGA CTT ACA GGT TAC CTC CGA AAC AAG TAA TTG TCT GAA TGT CCA ATG GAG GCT TTG.

A method of claim 52,7 wherein said DNA has the

## sequence:

ATG ACC AAC AAG TGT CTC CTC CAA ATT GCT CTC CTG TTG TGC TTC TCC TAC TGG TTG TTC ACA GAG GAG GAT TAA CGA GAG GAC AAC ACG AAG AGG ACT ACA GCT CTT TCC ATG AGC TAC AAC TTG CTT GGA TTC CTA CAA AGA TGA TGT CGA GAA AGG TAC TCG ATG TTG AAC GAA CCT AAG ATT GTT TCT AGC AGC AAT TTT CAG TGT CAG AAG CTC CTG TGG CAA TTG AAT GGG AGG TCG TCG TTA AAA GTC ACA GTC TTC GAG GAC ACC GTT AAC TTA CCC TCC CTT GAA TAT TGC CTC AAG GAC AGG ATG AAC TTT GAC ATC CCT GAG GAG GAA CTT ATA ACG GAG TTC CTG TCC TAC TTG AAA CTG TAG GGA CTC CTC ATT AAG CAG CAG CTC CTC

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58. Recombinant human fibroblast  $\beta_1$  interferon having the amino acid sequence:

Met Ser Tyr Asn Leu Leu Gly Phe Leu Gln Arg Ser Ser Asn Phe Gln Cys Gln Lys Leu Leu Trp Gln Leu Asn Gly Arg Leu Glu Tyr Cys Leu Lys Asp Arg Met Asn Phe Asp Ile Pro Glu Glu Ile Lys Gln Leu Gln Gln Phe Gln Lys Glu Asp Ala Ala Leu Thr Ile Tyr Glu Met Leu Gln Asn Ile Phe Ala Ile Phe Arg Gln Asp Ser Ser Ser Thr Gly Trp Asn Glu Thr Ile Val Elu Asn Leu Leu Ala Asn Val Tyr His Gln Ile Asn His Leu Lys Thr Val Leu Glu Glu Lys Leu Glu Lys Glu Asp Phe Thr Arg Gly Lys Leu Met Ser Ser Leu His Leu Lys Arg Tyr Tyr Gly Arg Ile Leu His Tyr Leu Lys Ala Lys Glu Tyr Ser His Cys Ala Trp Thr Ile Val Arg Val Glu Ile Leu Arg Asn Phe Tyr Phe Ile Asn Arg Leu Thr Gly Tyr Leu Arg Asn.

59. Recombinant human fibroblast  $eta_1$  interferon having the amino acid sequence:

Met Thr Asn Lys Cys Leu Leu Gln Ile Ala Leu Leu Leu Cys Phe Ser Thr Thr Ala Leu Ser Met Ser Tyr Asn Leu Leu Gly Phe Leu Gln Arg Ser Ser Asn Phe Gln Cys Gln Lys Leu Leu Trp Gln Leu Asn Gly Arg Leu Glu Tyr Cys Len Lys Asp Arg Met Asn Phe Asp Ile Pro Glu Glu Ile Lys Gln Leu Gln Asn Ile Phe Gln Lys Glu Asp Ala Ala Leu Thr Ile Tyr Glu Met Leu Gln Asn Ile Phe Ala Ile Phe Arg Gln Asp Ser Ser Ser Thr Gly Trp Asn Glu Thr Ile Val Glu Asn Leu Glu Asn Val Tyr His Gln Ile Asn His Leu Lys Thr Val Leu Glu Glu Lys Leu Glu Lys Glu Asp Phe Thr Arg Gly Lys Leu Met Ser Ser Leu His Leu Lys Arg Tyr Tyr Gly Arg Ile Leu His Tyr Leu Lys Ala Lys Glu Tyr Ser His Cys Ala Trp Thr Ile Val Arg Val Glu Ile Leu Arg Asn Phe Tyr Phe Ile Asn Arg Leu Thr Gly Tyr Leu Arg Asn. --

REMARKS

The specification has been amended to conform to the amendments made in Applicants' parent application Serial No. 06/201,359. The claims have also been amended to define Applicants' invention with the particularity required by statute and to set forth the invention in such manner as to clearly demonstrate its patentability.

Additionally, new claims 49-59 are submitted herewith to